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 421 aaaataccat ttttaaggcc tgggctgtag agacagggaa gtaccaggaa ggggtggatg
 481 accctgaccc agctaaatgg aaggcccagc tgcgctgtgc tctcaataag agcagagaat
 541 tcaacctgat gtatgatggc accaaggagg tgcccatgaa cccagtgaag atatatcaag
 601 tgtgtgacat ccctcagccc cagggctcga tcattaaccc aggatccaca gggtctgctc
 661 cctgggatga gaaggataat gatgtggatg aagaagatga ggaagatgag ctggatcagt
 721 cgcagcacca tgttcccatc caggacacct tccccttcct gaacatcaat ggttctccca
 781 tggcgccagc cagtgtgggc aattgcagtg tgggcaactg cagcccggag gcagtgtggc
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 961 gtgggaagga gtacgggcag accatgaccg tgagcaaccc tcagggctgc cgactcttct
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1321 gcgatctcat tgcccaccag aaaggacaga tagagaagca gccaccgttt gagatctact
1381 tatgctttgg ggaagaatgg ccagatggga aaccattgga aaggaaactc atcttggttc
1441 aggtcattcc agtagtggct cggatgatct acgagatgtt ttctggtgat ttcacacgat
1501 cctttgatag tggcagtgtc cgcctgcaga tctcaacccc agacatcaag gataacatcg
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1921 tggacaccaa agttctagct gcagctttgc ttccaatgtg accttgaaca agtcctttaa
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2041 catccagctt tagaacggac tcaatgacct tcttctactt gtacaaggct aaactgcctg
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2161 ctcaagcact a
```

Fig. 1

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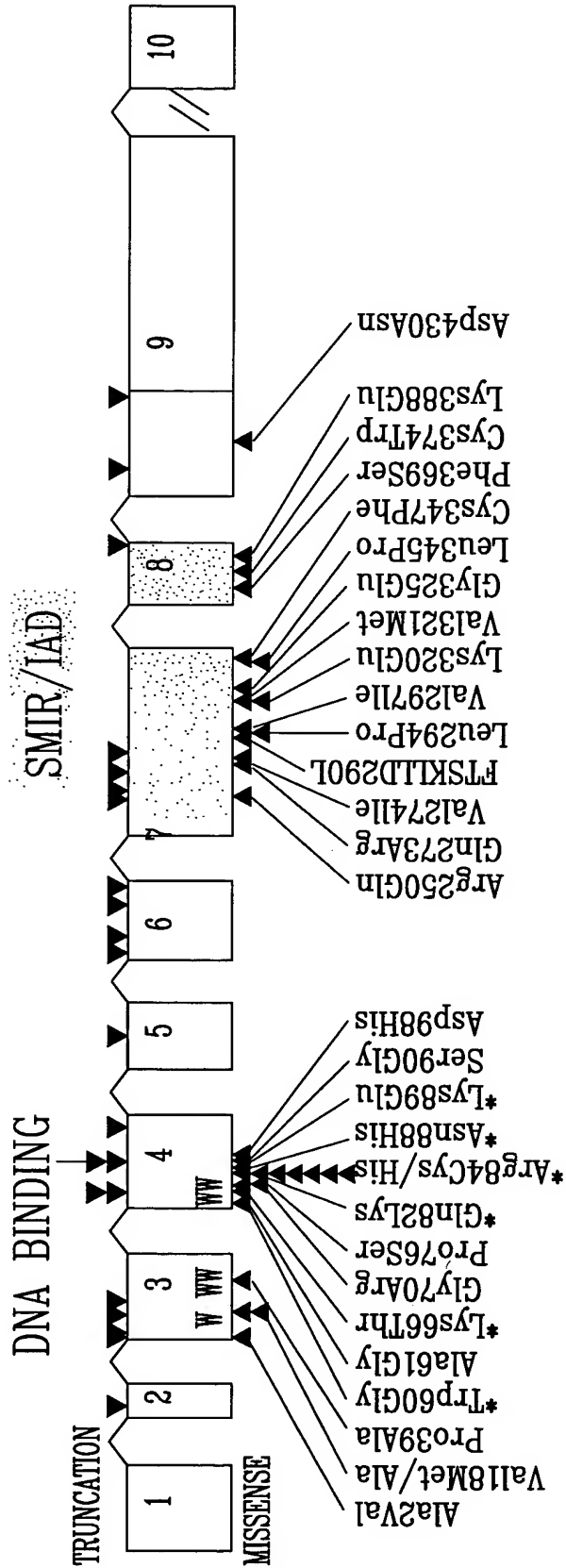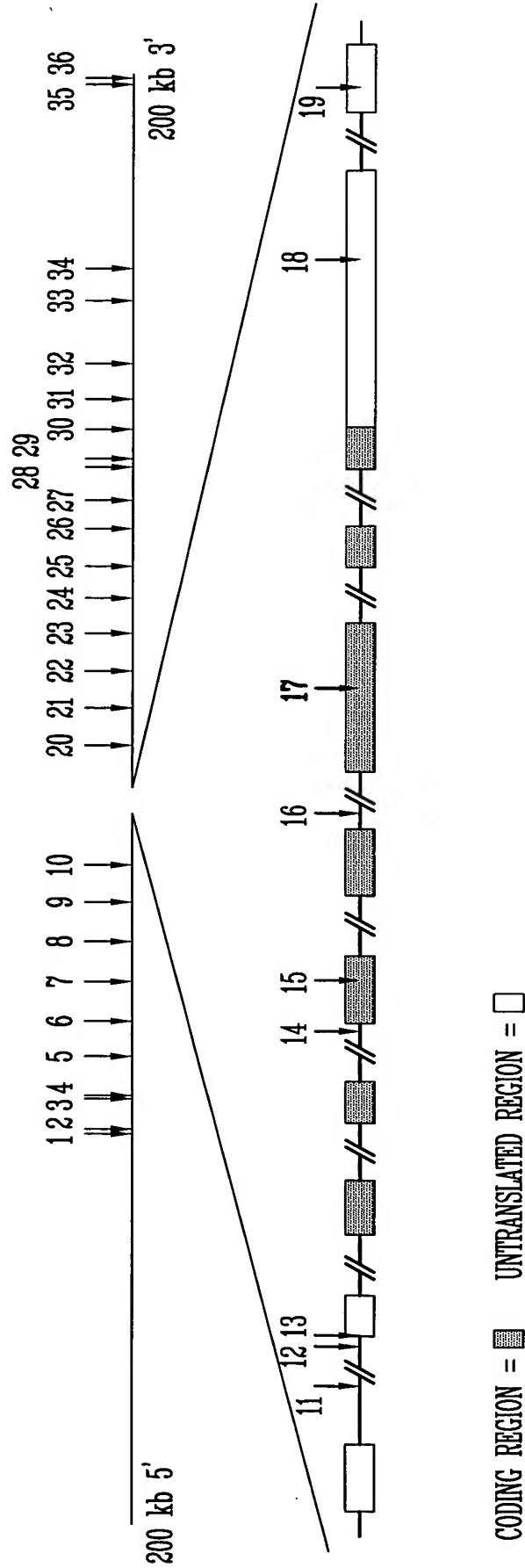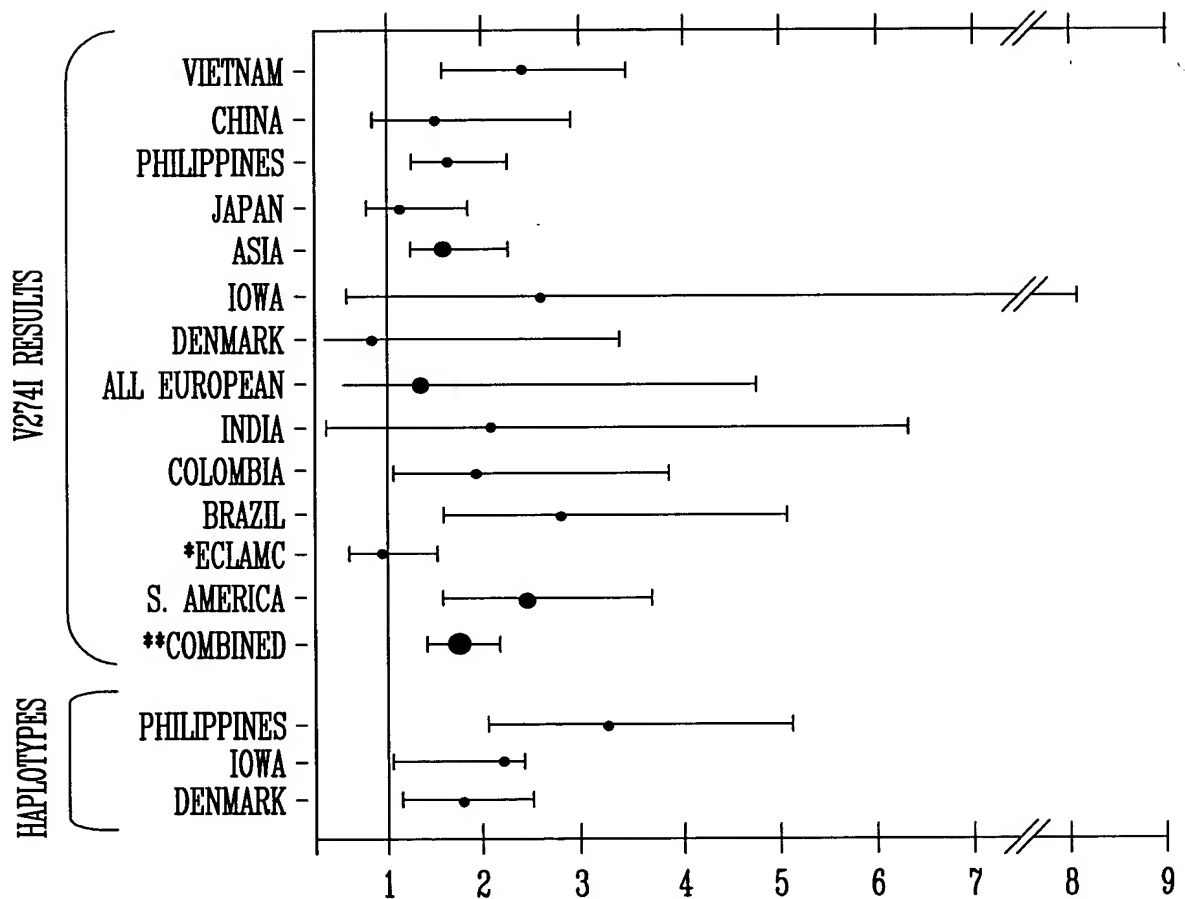QVIPVVARMIYEMFSGDFTRSFDSGSVRLQISTPDIKDNIVAQLKQLYRILQTQESWQ
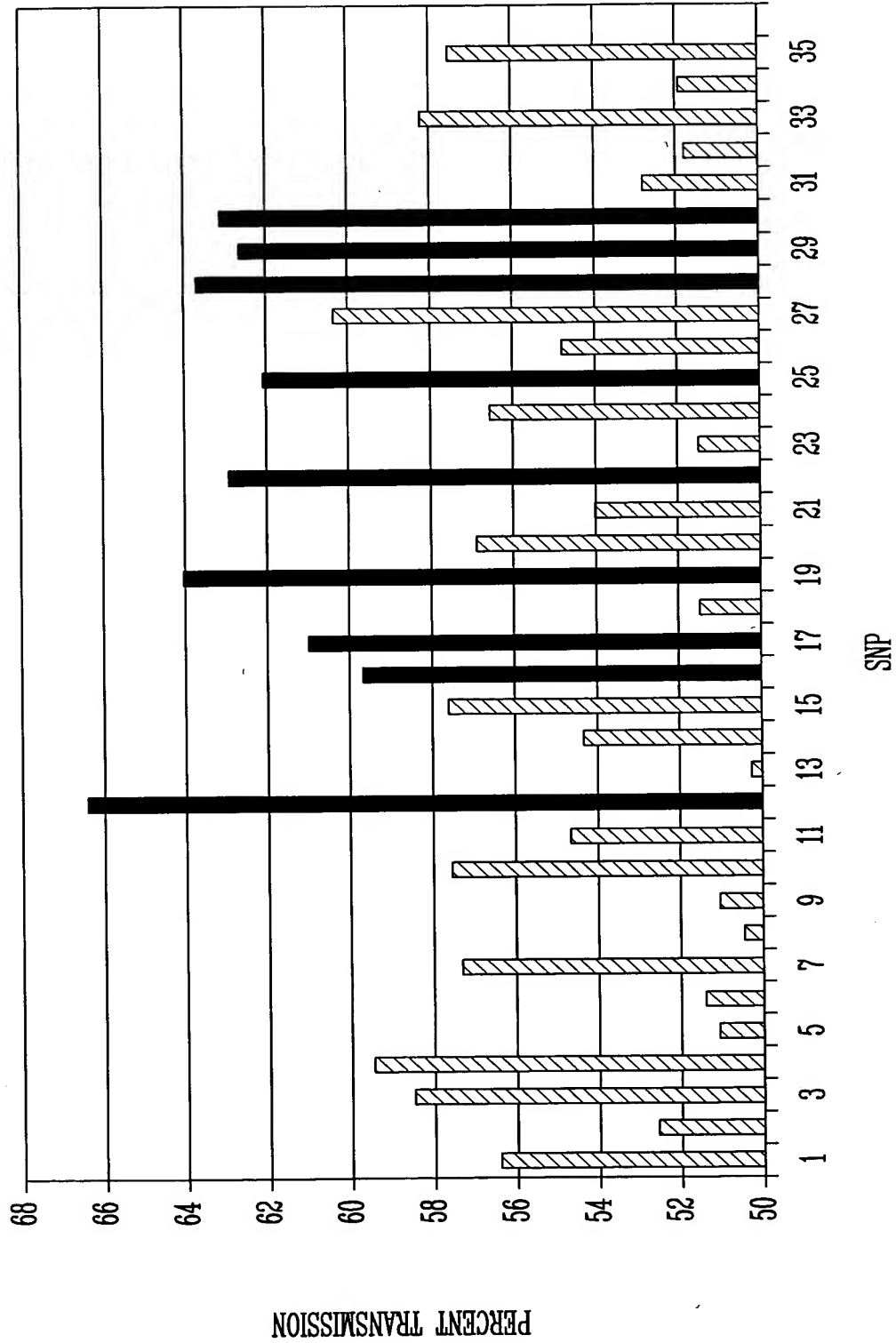PMQPTPSMQLPPALPPQ

Fig.2

Fig. 3
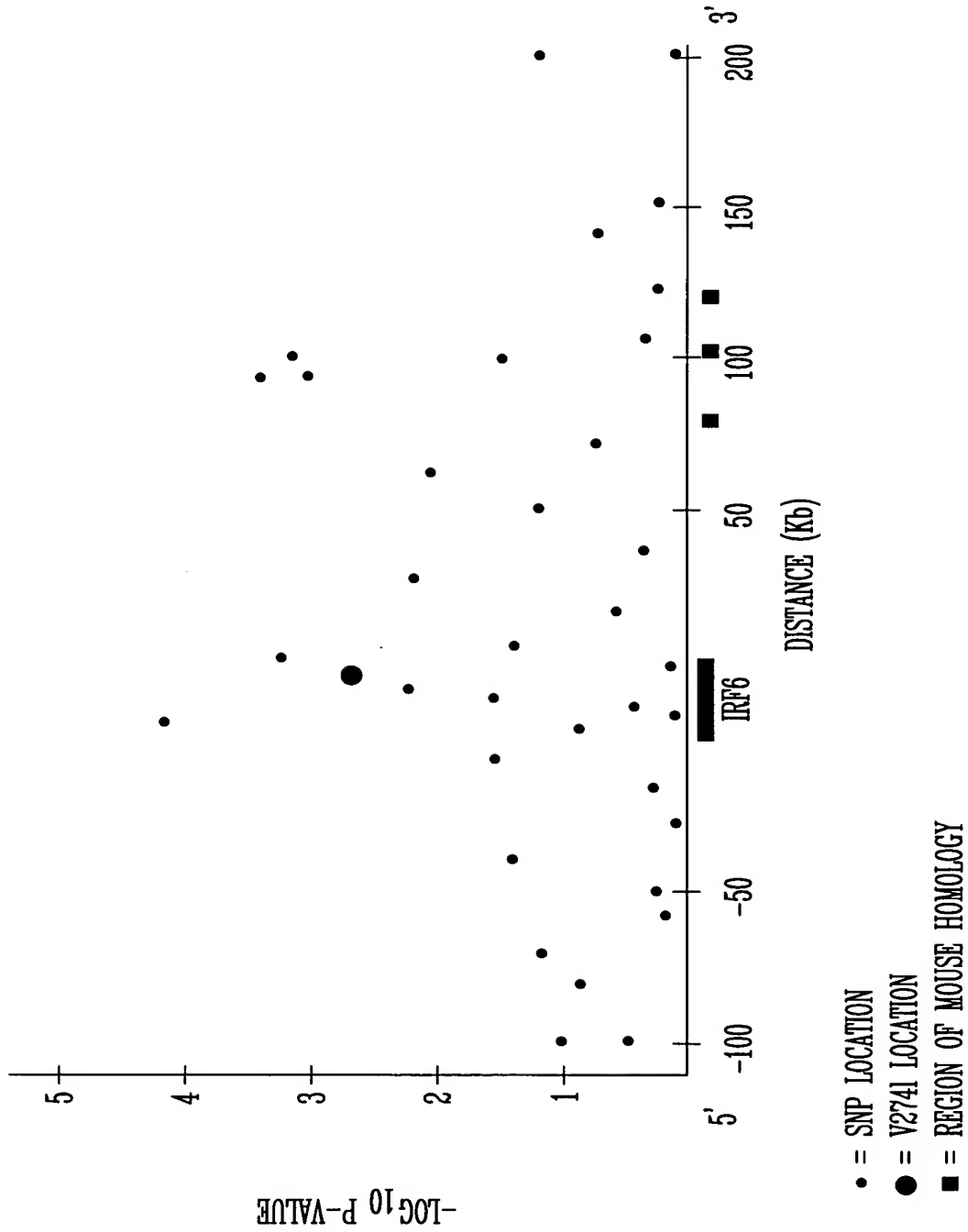
Fig. 4

CODING REGION = ▨   UNTRANSLATED REGION = ☐

*ECLAMC NOT INCLUDED IN COMBINED TOTAL
**COMBINED TOTAL IS OR FOR ALL V274I RESULTS

Fig. 5

Fig. 6

Fig. 7

HAPLOTYPE FOR SNP'S WITH FBAT P-VALUE <0.009

HAPLOTYPE	SNP 12	SNP 16	SNP 17	SNP 19	SNP 22	SNP 25	SNP 27	SNP 28	SNP 29	FREQUENCY	P-VALUE
1										0.46	0.0002 (+)
2										0.08	0.07
3										0.07	0.07
4										0.07	0.08
5										0.04	0.24
OTHERS										0.28	

HAPLOTYPE >1% IN DANISH AND IOWA TRIADS

HAPLOTYPE	SNP 1	SNP 6	SNP 8	SNP 16	FREQUENCY	P-VALUE
1					0.54	<0.001 (+)
2					0.19	0.03 (−)
3					0.12	0.09
4					0.09	0.77
OTHERS					0.06	

COMMON ALLELE AT SNP = ■   RARE ALLELE AT SNP = □

Fig. 8